Hit List

Clear Generate Collection Print Fwd Refs Bkwd Refs
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Search Results - Record(s) 1 through 5 of 5 returned.

1. Document ID: US 6787133 B2

L2: Entry 1 of 5

File: USPT

Sep 7, 2004

US-PAT-NO: 6787133

DOCUMENT-IDENTIFIER: US 6787133 B2

TITLE: Using purified telomerase to identify telomerase activators and inhibitors

DATE-ISSUED: September 7, 2004

INVENTOR-INFORMATION:

NAME CITY STATE ZIP CODE COUNTRY

Weinrich; Scott L. Chesterfield MO
Atkinson, III; Edward M. Seattle WA
Lichtsteiner; Serge P. Encinitas CA
Vasserot; Alain P. Berkeley CA

Pruzan; Ronald A. Palo Alto CA

US-CL-CURRENT: $\underline{424}/\underline{94.5}$; $\underline{435}/\underline{14}$, $\underline{435}/\underline{194}$, $\underline{435}/\underline{252.3}$, $\underline{435}/\underline{320.1}$, $\underline{435}/\underline{412}$, $\underline{435}/\underline{413}$, $\underline{435}/\underline{6}$, $\underline{435}/\underline{8}$, $\underline{435}/\underline{9}$, $\underline{435}/\underline{91.3}$, $\underline{435}/\underline{935}$, $\underline{530}/\underline{350}$, $\underline{536}/\underline{23.2}$

Full Title Citation Front Review Classification Date Reference

2. Document ID: US 6545133 B1

L2: Entry 2 of 5

File: USPT

Apr 8, 2003

US-PAT-NO: 6545133

DOCUMENT-IDENTIFIER: US 6545133 B1

TITLE: Methods for purifying telomerase

DATE-ISSUED: April 8, 2003

INVENTOR-INFORMATION:

NAME CITY STATE ZIP CODE COUNTRY

Weinrich; Scott L. Chesterfield MO
Atkinson, III; Edward M. Seattle WA
Lichtsteiner; Serge P. Encinitas CA
Vasserot; Alain P. Berkeley CA

Page 2 of 3 Record List Display

Pruzan; Ronald A.

Palo Alto CA

US-CL-CURRENT: 530/413; 435/194, 435/252.3, 435/320.1, 530/355, 530/358, 536/23.2,

<u>536/23.5</u>, <u>536/24.31</u>

Full Title Citation Front Review Classification Date Reference Citation Claims KMC Draw De

3. Document ID: US 6517834 B1

L2: Entry 3 of 5

File: USPT

Feb 11, 2003.

US-PAT-NO: 6517834

DOCUMENT-IDENTIFIER: US 6517834 B1

TITLE: Purified telomerase

DATE-ISSUED: February 11, 2003

INVENTOR-INFORMATION:

NAME CITY STATE ZIP CODE COUNTRY

Weinrich; Scott L. Chesterfield MO Atkinson, III; Edward M. WA Seattle Lichtsteiner; Serge P. Encinitas CA Vasserot; Alain P. CA Berkeley Pruzan; Ronald A. Palo Alto CA

US-CL-CURRENT: 424/94.5; 435/194, 435/252.3, 435/320.1, 435/91.3, 530/412, 530/413,

536/23.2

Full Title Citation Front Review Classification Date Reference

4. Document ID: US 6261556 B1

L2: Entry 4 of 5

File: USPT

Jul 17, 2001

US-PAT-NO: 6261556

DOCUMENT-IDENTIFIER: US 6261556 B1

** See image for <u>Certificate of Correction</u> **

TITLE: Purified telomerose

DATE-ISSUED: July 17, 2001

INVENTOR-INFORMATION:

NAME CITY STATE ZIP CODE COUNTRY

Weinrich; Scott L. Redwood City CA Atkinson, III; Edward M. Seattle WA Lichtsteiner; Serge P. Cupertino CA Vasserot; Alain P. Saratoga CA Pruzan; Ronald A. Palo Alto CA

Record List Display Page 3 of 3

Kealey; James T.

San Anselmo

CA

US-CL-CURRENT: 424/94.5; 435/194, 435/252.3, 435/320.1, 435/91.3, 530/412, 530/413, 536/23.2

Full	Title	Citation Front	Review	Classification	Date	Reference		Claims	KWIC	Draw De
m	5	Document ID:	115 506	58506 Δ	••••		***************************************			

5. Document ID: US 5968506 A

L2: Entry 5 of 5

File: USPT

Oct 19, 1999

US-PAT-NO: 5968506

DOCUMENT-IDENTIFIER: US 5968506 A

TITLE: Purified telomerase

DATE-ISSUED: October 19, 1999

INVENTOR-INFORMATION:

NAME CITY STATE ZIP CODE COUNTRY Weinrich; Scott L. Redwood City CA Atkinson, III; Edward M. Seattle WA Lichtsteiner; Serge P. CA Cupertino Vasserot; Alain P. Saratoga CA Pruzan; Ronald A. Palo Alto CA Kealey; James T. San Anselmo CA

US-CL-CURRENT: 424/94.5; 435/194, 435/252.3, 435/320.1, 435/91.3, 530/412, 530/413, 536/23.2

Full Title Citation Front	Review Classification Date	Reference	Claims KNAC Draw D
Clear Generate Col	lection Print F	wd Refs Bkwd Refs	Generate OACS
Terms		Documents	·
L1 and 2000-fold			5

Display Format: CIT Change Format

Previous Page Next Page Go to Doc#

Record Display Form Page 1 of 3

First Hit Fwd Refs Previous Doc Next Doc Go to Doc# Generate Collection Print

L2: Entry 1 of 5 File: USPT Sep 7, 2004

US-PAT-NO: 6787133

DOCUMENT-IDENTIFIER: US 6787133 B2

TITLE: Using purified telomerase to identify telomerase activators and inhibitors

DATE-ISSUED: September 7, 2004

INVENTOR-INFORMATION:

NAME	CITY	STATE	ZIP CODE	COUNTRY
Weinrich; Scott L.	Chesterfield	MO		
Atkinson, III; Edward M.	Seattle	WA		
Lichtsteiner; Serge P.	Encinitas	CA		
Vasserot; Alain P.	Berkeley	CA		
Pruzan; Ronald A.	Palo Alto	CA		

US-CL-CURRENT: 424/94.5; 435/14, 435/194, 435/252.3, 435/320.1, 435/412, 435/413, 435/6, 435/8, 435/9, 435/91.3, 435/935, 530/350, 536/23.2

CLAIMS:

The invention claimed is:

- 1. A method to identify a regulator of telomerase activity, comprising: a) obtaining a preparation of mammalian telomerase enzyme purified by at least <u>2000-fold</u> more pure than an extract of cells from adenovirus-transformed kidney cell line (293 cells), wherein the telomerase enzyme contains telomerase RNA component, and has a molecular weight of 200-2000 kDa; b) combining the preparation with a compound; c) determining telomerase activity of the enzyme in the presence of the compound; d) identifying the compound as being a regulator of telomerase if the telomerase activity measured in step c) is affected by the presence of the compound.
- 2. The method of claim 1, wherein the telomerase preparation was obtained by a process in which a solution containing telomerase activity was combined with an oligonucleotide having specific activity for mammalian telomerase; and then protein was collected that had bound the oligonucleotide.
- 3. The method of claim 2, wherein the oligonucleotide comprises a retrievable label.
- 4. The method of claim 3, wherein the retrievable label is biotin.
- 5. The method of claim 2, wherein the solution that was combined with the oligonucleotide had been obtained by preparing an enriched solution from a cell expressing telomerase, whereby telomerase enzyme in the enriched solution was separated from other proteins expressed by the cell.

- 6. The method of claim 2, wherein the process used to prepare the telomerase comprised combining a fraction containing telomerase enzyme with an anion exchange matrix, and collecting protein that bound the matrix.
- 7. The method of claim 2, wherein the process used to prepare the telomerase comprised combining a fraction containing telomerase enzyme with a cation exchange matrix (such as a heparin matrix), and collecting protein that bound the matrix.
- 8. The method of claim 2, wherein the process used to prepare the telomerase comprised combining a fraction containing telomerase enzyme with an intermediate selectivity matrix, and collecting protein that bound the matrix; wherein the intermediate selectivity matrix had at least one of the following substituents: hydroxyapatite, a polyamine (such as spermine or spermidine), poly guanylic acid, a divalent metal ion (such as Ni.sup.++), a positively charged poly-amino acid (such as poly-L-lysine), a positively charged enzyme (such as histone), or aminophenyl-boronic acid.
- 9. The method of claim 2, wherein the process used to prepare the telomerase comprised separating a fraction containing the telomerase enzyme by gel filtration chromatography or gradient centrifugation that separates molecules $>200\ kDa$.
- 10. The method of claim 2, wherein the oligonucleotide contains a sequence that binds specifically to telomerase RNA component.
- 11. The method of claim 10, wherein the oligonucleotide contains the sequence of oligo 5 (SEQ. ID NO:3).
- 12. The method of claim 2, wherein the oligonucleotide contains a sequence that is specifically recognized by telomerase protein.
- 13. The method of claim 12, wherein the oligonucleotide contains the sequence (TTAGGG).sub.3 (SEQ. ID NO:6).
- 14. The method of claim 12, wherein the oligonucleotide does not contain the sequence (TTAGGG).sub.3 (SEQ. ID NO:6).
- 15. The method of claim 12, wherein the oligonucleotide contains the sequence of M2/TS (SEQ. ID N0:8).
- 16. The method of claim 1, wherein the telomerase preparation is at least .about.20,000 fold more pure than the cell extract.
- 17. The method of claim 1, wherein the telomerase preparation is between .about.3,000 and .about.60,000 fold more pure than the cell extract.
- 18. The method of claim 1, wherein the telomerase protein is human.
- 19. The method of claim 1, wherein the telomerase preparation has measurable telomerase activity in 0.2 .mu.g of protein when quantified in a telomere primer elongation assay in which .sup.32 P-labeled primer extensions are separated on a gel and detected using a phosphoimager screen.
- 20. The method of claim 1, wherein telomerase core enzyme is present in the

- preparation at a concentration of at least 3.times.10.sup.-10 mol L.sup.-1.
- 21. The method of claim 1, wherein telomerase core enzyme is present in the preparation at a concentration of at least 2.times.10.sup.-9 mol L.sup.-1.
- 22. The method of claim 1, wherein the telomerase activity is determined in step c) by a primer elongation assay.
- 23. The method of claim 1, wherein the telomerase activity is determined in step c) by a dot blot assay.
- 24. The method of claim 1, whereby the compound is identified as being an inhibitor of telomerase.
- 25. The method of claim 1, whereby the compound is identified as being an activator of telomerase.

Previous Doc Next Doc Go to Doc#

WEST Search History

Hide Items Restore Clear Cancel

DATE: Monday, April 18, 2005

Hide?	Set Name	<u>e Query</u>	Hit Count
	DB=PG	PB,USPT,USOC,EPAB,JPAB,DWPI; PLUR=YES; OP=	ADJ
	L6	telomerase with activator with inhibitor.clm.	2
	L5	telomerase with activator with inhibitor with 2000-fold	0
	L4	telomerase with activator with inhibitor	60
	L3	telomerase same activator same inhibitor	209
	DB = US	PT; PLUR=YES; OP=ADJ	
	L2	L1 and 2000-fold	5
	L1	telomerase same purification	66

END OF SEARCH HISTORY

FILE 'HOME' ENTERED AT 10:49:44 ON 18 APR 2005

=> file medline hcaplus scisearch embase

COST IN U.S. DOLLARS SINCE FILE TOTAL ENTRY SESSION

FULL ESTIMATED COST ENTRY SESSION 0.84 0.84

FILE 'MEDLINE' ENTERED AT 10:51:57 ON 18 APR 2005

FILE 'HCAPLUS' ENTERED AT 10:51:57 ON 18 APR 2005
USE IS SUBJECT TO THE TERMS OF YOUR STN CUSTOMER AGREEMENT.
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=> s mammalian telomerase and (kidney cell line or 293 cell?)
L1 4 MAMMALIAN TELOMERASE AND (KIDNEY CELL LINE OR 293 CELL?)

=> dup rem 11

PROCESSING COMPLETED FOR L1

L2 4 DUP REM L1 (0 DUPLICATES REMOVED)

=> d 12 1-4 ibib ab

L2 ANSWER 1 OF 4 HCAPLUS COPYRIGHT 2005 ACS on STN

ACCESSION NUMBER:

2003:276752 HCAPLUS Full-text

DOCUMENT NUMBER:

138:283318

TITLE:

Sequential purification of mammalian

telomerase including affinity chromatography

using oligonucleotide sorbents

INVENTOR(S):

Weinrich, Scott L.; Atkinson, Edward M., III;

Lichtsteiner, Serge P.; Vasserot, Alain P.; Pruzan,

Ronald A.

PATENT ASSIGNEE(S):

Geron Corporation, USA

SOURCE:

U.S., 24 pp., Cont.-in-part of U.S. 6,261,556.

CODEN: USXXAM

DOCUMENT TYPE:

Patent

LANGUAGE:

English

FAMILY ACC. NUM. COUNT:

3

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
US 6545133	B1	20030408	US 2000-717829	20001120
US 5968506	Α	19991019	US 1997-833377	19970404
US 6261556	B1	20010717	US 1999-420056	19991018
PRIORITY APPLN. INFO.:			US 1995-510736 B	2 19950804
			US 1997-833377 A	1 19970404

US 1999-420056 A2 19991018

This invention provides purified mammalian telomerase and methods of AB purifying it. The methods involve the use of several sequential steps, including the use of anion exchange matrix, heparin-containing matrix, spermidine-containing matrixes, gel filtration chromatog. or gradient centrifugation, and affinity purification An affinity agent (oligonucleotide complementary to the RNA component of telomerase labeled with biotin and isolated with matrix-bound streptavidin) is disclosed. A method for preparing human telomerase that is 65,320-fold purified compared to that in crude cell extract is described. The method comprises six steps in succession: (1) CHAPS detergent S-100 extract preparation from 293 cells; (2) chromatog. of the S-100 extract on POROS 50HQ matrix; (3) chromatog. of the POROS 50HQ active fractions on POROS Heparin 20HE-1 matrix; (4) chromatog. of the POROS Heparin 20 HE-1 active fractions on POROS spermidine matrix; (5) chromatog. of the POROS Spermidine active fractions on Superose 6 sizing column; and (6) chromatog. of the Superose 6 sizing column active fractions on Oligo 5 affinity matrix.

REFERENCE COUNT: 43 THERE ARE 43 CITED REFERENCES AVAILABLE FOR

THIS

RECORD. ALL CITATIONS AVAILABLE IN THE RE

FORMAT

L2 ANSWER 2 OF 4 HCAPLUS COPYRIGHT 2005 ACS on STN ACCESSION NUMBER: 2003:113309 HCAPLUS Full-text

DOCUMENT NUMBER: 138:165719

TITLE: Chromatographic purification of human telomerase

from

293 cells

INVENTOR(S): Weinrich, Scott L.; Atkinson, Edward M., III;

Lichtsteiner, Serge P.; Vasserot, Alain P.; Pruzan,

Ronald A.

PATENT ASSIGNEE(S): Geron Corporation, USA

SOURCE: U.S., 24 pp., Cont.-in-part of U.S. 6,261,556.

CODEN: USXXAM

DOCUMENT TYPE: Patent LANGUAGE: English

FAMILY ACC. NUM. COUNT: 3

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
US 6517834	B1	20030211	US 2000-717828	20001120
US 5968506	A	19991019	US 1997-833377	19970404
US 6261556	B1	20010717	US 1999-420056	19991018
US 2003186282	A1	20031002	US 2002-330872	20021224
US 6787133	B2	20040907		
PRIORITY APPLN. INFO.:			US 1995-510736	B2 19950804
			US 1997-833377	Al 19970404
			US 1999-420056	A2 19991018
			US 2000-717828	A1 20001120

AB This invention provides purified human telomerase and methods of purifying it. The methods involve the use of several sequential steps, including the use of matrixes that bind mols. bearing neg. charges, matrixes that bind mols. bearing pos. charges, intermediate-selectivity matrixes, methods that sep. mols. based on their size, shape, or buoyant d., and by affinity purification Human telomerase was purified to over

60,000-fold purity from 293 cell exts. This method comprises six steps in succession: (1) CHAPS detergent S-100 extract preparation from 293 cells; (2) chromatog. of the S-100 extract on POROS 50HQ matrix; (3) chromatog. of the POROS 50HQ active fractions of POROS Heparin 20HE1 matrix; (4) chromatog. of the POROS Heparin 20HE1 active fractions on POROS spermidine matrix; (5) chromatog. of the POROS Spermidine active fractions on Superose 6 sizing column; and (6) chromatog. of the Superose 6 sizing column active fractions on Oligo 5 affinity matrix. A telomere primer elongation assay for mammalian telomerase is also described.

REFERENCE COUNT:

45 THERE ARE 45 CITED REFERENCES AVAILABLE FOR

THIS

RECORD. ALL CITATIONS AVAILABLE IN THE RE

FORMAT

L2 ANSWER 3 OF 4 SCISEARCH COPYRIGHT (c) 2005 The Thomson Corporation on

STN

ACCESSION NUMBER:

2001:695761 SCISEARCH Full-text

THE GENUINE ARTICLE: 466BD

TITLE:

Human telomerase RNA-protein interactions

AUTHOR: CORPORATE SOURCE: Bachand F; Triki F; Autexier C (Reprint)
Sir Mortimer B Davis Jewish Hosp, Lady Davis Inst Med

Res,

Bloomfield Ctr Res Aging, 3755 Cote St Catherine Rd, Montreal, PQ H3T 1E2, Canada (Reprint); Sir Mortimer B Davis Jewish Hosp, Lady Davis Inst Med Res, Bloomfield

Ctr

Res Aging, Montreal, PQ H3T 1E2, Canada; McGill Univ,

Dept

Anat & Cell Biol, Montreal, PQ H3A 2B2, Canada

COUNTRY OF AUTHOR:

Canada

SOURCE:

NUCLEIC ACIDS RESEARCH, (15 AUG 2001) Vol. 29, No. 16,

pp.

3385-3393.

Publisher: OXFORD UNIV PRESS, GREAT CLARENDON ST, OXFORD

OX2 6DP, ENGLAND. ISSN: 0305-1048. Article; Journal

DOCUMENT TYPE:

LANGUAGE: English

REFERENCE COUNT:

42

ABSTRACT IS AVAILABLE IN THE ALL AND IALL FORMATS

AB Telomere length is maintained in most eukaryotic cells by telomerase. The core components of this ribonucleoprotein (RNP) enzyme include a protein catalytic subunit, composed of motifs conserved among reverse transcriptases (RT), and an RNA subunit that contains a short template sequence essential for the synthesis of telomeric repeats. We developed an electrophoretic mobility shift assay using active telomerase partially purified from 293 cells and radiolabeled, in vitro-transcribed human telomerase RNA (hTR) to investigate the molecular interactions of the human telomerase RT (hTERT) and telomerase-associated proteins with hTR. A specific hTR-protein complex was identified and shown to contain hTERT and human Staufen by antibody supershift assays. Variants of hTR altered in distinct structural elements were analyzed for their ability to competitively inhibit complex formation. Human telomerase RNAs lacking the CR4-CR5 domain were poor inhibitors of hTR-protein complex formation, suggesting that the CR4-CR5 domain of hTR is a potential proteinbinding site. Furthermore, alterations in the telomerase RNA pseudoknot's P3 helix, the CR7 domain, or the H/ACA box efficiently inhibited formation of the complex, indicating that these domains are dispensable for the assembly of a telomerase RNP in vitro. Potential telomerase-associated proteins that bind hTR were also identified using a UV cross-linking assay.

L2ANSWER 4 OF 4 SCISEARCH COPYRIGHT (c) 2005 The Thomson Corporation on

ACCESSION NUMBER: 1999:410137 SCISEARCH Full-text

THE GENUINE ARTICLE: 199DA

TITLE: Genomic organization and promoter characterization of

the

gene encoding the human telomerase reverse transcriptase

(hTERT)

AUTHOR: Wick M (Reprint); Zubov D; Hagen G

BAYER AG, DIV CENT RES, DEPT MOL BIOL, D-51368 CORPORATE SOURCE:

LEVERKUSEN,

GERMANY (Reprint)

COUNTRY OF AUTHOR: **GERMANY**

SOURCE:

GENE, (17 MAY 1999) Vol. 232, No. 1, pp. 97-106. Publisher: ELSEVIER SCIENCE BV, PO BOX 211, 1000 AE

AMSTERDAM, NETHERLANDS.

ISSN: 0378-1119.

DOCUMENT TYPE:

Article; Journal

FILE SEGMENT: LANGUAGE:

LIFE English

REFERENCE COUNT:

ABSTRACT IS AVAILABLE IN THE ALL AND IALL FORMATS

AB The enzyme telomerase plays a crucial role in cellular proliferation and tumorigenesis. By adding hexameric repeats to chromosome ends, it prevents telomeric loss and, thus, entry into senescence. Recent data suggest that expression of the human telomerase reverse transcriptase subunit (hTERT) represents the limiting factor for telomerase activity. To gain an insight into the mechanisms regulating hTERT expression, we have determined the complete genomic organization of the hTERT gene and isolated the 5'- and 3'- flanking region. The hTERT gene encompasses more than 37 kb and consists of 16 exons. We show that all hTERT insertion and deletion variants described so far most likely result from the usage of alternative splice consensus sequences in intron or exon regions. Furthermore, we identified a new hTERT splice variant. Analysis of the DNA sequence surrounding the putative transcriptional start region revealed a TATA-less promoter located in a CpG island. A promoter fragment spanning the first 1100 bp upstream of the initiating ATG start codon exhibited high-level activity in HEK-293 cells. Several consensus binding sites for the transcription factor Spl as well as a c-Myc binding site were identified in this promoter region. Altogether, these results provide the basis for more detailed studies on the regulation of telomerase activity in normal and cancer cells, and may lead to the development of new cancer therapies. (C) 1999 Elsevier Science B.V. All rights reserved.

```
=> s mammalian telomerase and (inhibitor? or activator? or regulator?)
           47 MAMMALIAN TELOMERASE AND (INHIBITOR? OR ACTIVATOR? OR
L3
REGULATOR?
              )
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PROCESSING COMPLETED FOR L3
            47 DUP REM L3 (0 DUPLICATES REMOVED)
=> focus 14
PROCESSING COMPLETED FOR L4
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=> d 15 1-10 ibib ab
    ANSWER 1 OF 47 HCAPLUS COPYRIGHT 2005 ACS on STN
ACCESSION NUMBER: 1998:184000 HCAPLUS Full-text
                      128:240310
DOCUMENT NUMBER:
TITLE:
                       Chimeric telomerase gene promoter-reporter gene
assays
                       for regulators of mammalian
                       telomerase expression
INVENTOR(S):
                       Villeponteau, Bryant; Harley, Calvin
PATENT ASSIGNEE(S): Geron Corporation, USA; Villeponteau, Bryant;
Harley,
                       Calvin
                       PCT Int. Appl., 59 pp.
SOURCE:
                       CODEN: PIXXD2
DOCUMENT TYPE:
                       Patent
LANGUAGE:
                       English
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
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                                                             DATE
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    WO 9811207
                       A2
                             19980319
                                                            19970916
    WO 9811207
WO 9811207
                                         WO 1997-US16450
                      A3
                             19980625
        W: AU, CA, CN, JP, KR, MX, US
        RW: AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT,
SE
    US 5972605
AU 9743519
                      Α
                             19991026 US 1996-714482
                                                              19960916
                       A1 19980402 AU 1997-43519
                                                              19970916
                                                          A1 19960916
PRIORITY APPLN. INFO.:
                                         US 1996-714482
                                                           B2 19940707
                                         US 1994-272102
                                         US 1994-330123
                                                           A2 19941027
                                         US 1995-472802
                                                          A2 19950607
                                         US 1995-482115
                                                          A2 19950607
                                         US 1995-521634
                                                           B2 19950831
                                         WO 1997-US16450
                                                          W 19970916
     Telomerase reporter constructs are suitable for use in reporting
     transcriptional activity of a mammalian telomerase gene transcription
     regulatory region. The constructs contain a transcription regulatory
     region of a mammalian telomerase gene operably linked to a reporter
    polynucleotide sequence. The regulatory region for the gene encoding
    the RNA component of human telomerase is reported and may be used in
```

constructs containing such reporter genes as CAT, β -GAL, NEOR, HSV-TK to

create recombinant mammalian host cells that can used to identify

telomerase transcription modulators, which are potential anticarcinogenic agents.

ANSWER 2 OF 47 HCAPLUS COPYRIGHT 2005 ACS on STN

ACCESSION NUMBER:

1999:686603 HCAPLUS Full-text

DOCUMENT NUMBER:

131:318570

TITLE:

Telomerase reporter constructs suitable for use in

reporting activity of the transcription

regulatory region of a mammalian

telomerase gene

INVENTOR(S):

Villeponteau, Bryant; Harley, Calvin

PATENT ASSIGNEE(S):

Geron Corporation, USA

SOURCE:

U.S., 22 pp., Cont.-in-part of U.S. Ser. No.

521,634,

abandoned.

CODEN: USXXAM

DOCUMENT TYPE:

Patent

LANGUAGE:

English

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
US 5972605	Α	19991026	US 1996-714482	19960916
US 5583016	Α	19961210	US 1994-330123	19941027
US 5776679	Α	19980707	US 1995-482115	19950607
US 5958680	A	19990928	US 1995-472802	19950607
WO 9811207	. A2	19980319	WO 1997-US16450	19970916
WO 9811207	A3	19980625		
W: AU, CA, CN,	JP, KR	, MX, US		•
RW: AT, BE, CH,	DE, DK	, ES, FI, FF	R, GB, GR, IE, IT,	LU, MC, NL, PT,
SE	•			
AU 9743519	A1	19980402	AU 1997-43519	19970916
PRIORITY APPLN. INFO.:			US 1994-272102	B2 19940707
•			US 1994-330123	A2 19941027
			US 1995-472802	A2 19950607
			US 1995-482115	A2 19950607
			US 1995-521634	B2 19950831
			US 1996-714482	A1 19960916
,			WO 1997-US16450	W 19970916

AΒ The invention provides telomerase reporter constructs suitable for use in reporting activity of the transcription regulatory region of a mammalian telomerase gene. Said constructs comprise a human telomerase gene transcription regulatory region operably linked to a reporter polynucleotide sequence. In one embodiment, the transcription regulatory region comprises sequences from the hTR promoter region. certain embodiments, the invention provides for the use of the disclosed reporter constructs in assays for determining whether an agent modulates expression of telomerase.

REFERENCE COUNT:

11 THERE ARE 11 CITED REFERENCES AVAILABLE FOR

THIS

RECORD. ALL CITATIONS AVAILABLE IN THE RE

FORMAT

ANSWER 3 OF 47 HCAPLUS COPYRIGHT 2005 ACS on STN ACCESSION NUMBER: 1997:684422 HCAPLUS Full-text DOCUMENT NUMBER:

128:1459

TITLE:

Inhibitor peptide nucleic acids binding the

RNA component of mammalian

telomerase

INVENTOR (S):

Shay, Jerry W.; Wright, Woodring E.; Piatyszek,

Mieczyslaw A.; Corey, David; Norton, James C.

PATENT ASSIGNEE(S):

Geron Corp., USA

SOURCE:

PCT Int. Appl., 75 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent

LANGUAGE:

English

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 9738013	A1	19971016	WO 1997-US5931	19970409

W: AU, CA, CN, JP, KR, MX

RW: AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT,

SE

20						
US 6015710	Α	20000118	US	1996-630019		19960409
AU 9726631	A1	19971029	AU	1997-26631		19970409
JP 2001517929	T2	20011009	JP	1997-536487		19970409
PRIORITY APPLN. INFO.:			US	1996-630019	Α	19960409
			WO	1997-US5931	M	19970409

AB Peptide nucleic acids (PNAs) that can bind with the RNA moiety of mammalian telomerases and that can inhibit the enzyme are described. The PNAs may be antisense or triple helix- or D-loop-forming. The PNAs may be further modified with lipid moieties or signal peptides to ensure their efficient uptake by animal cells. The PNAs can be used to assay telomerase activity and to inhibit the enzyme in the treatment of disease. A series of PNA candidates for inhibition of telomerase activity were tested for efficacy in a telomere repeat amplification protocol assay and inhibition in the micromolar or nanomolar range was found. Further optimization expts. are reported.

L5 ANSWER 4 OF 47 HCAPLUS COPYRIGHT 2005 ACS on STN

ACCESSION NUMBER:

2000:351105 HCAPLUS Full-text

DOCUMENT NUMBER:

133:146531

TITLE:

The NACHT family - a new group of predicted NTPases

implicated in apoptosis and MHC transcription

activation

AUTHOR (S):

Koonin, Eugene V.; Aravind, L.

CORPORATE SOURCE:

National Center for Biotechnology Information,

National Library of Medicine, National Institutes of

Health, Bethesda, MD, 20894, USA

SOURCE:

Trends in Biochemical Sciences (2000), 25(5), 223-

224

CODEN: TBSCDB; ISSN: 0376-5067

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DOCUMENT TYPE: Journal LANGUAGE: English

In the course of the recent anal. of the domain architectures of proteins involved in programmed cell death, we noticed that neuronal apoptosis inhibitor protein (NAIP) and MHC class II transcription activator (CIITA) contain a distinct predicted nucleoside triphosphatase

(NTPase) domain. Here we report that this domain belongs to a new family of predicted NTPases that include animal, fungal and bacterial proteins. The existence of a new family became apparent with the recent identification of a new pro-apoptotic protein, CARD4, which activates NF-kB. CARD4 contains a predicted NTPase domain that shows highly significant sequence similarity to CIITA. When the sequence of CARD4 between positions 119 and 417 was used as the query for searching the non-redundant protein database at the NCBI with the gapped BLAST program, the random expectation (E) value for the alignment with CIITA was <10-14. A single iteration of this database search using the PSI-BLAST program (with the cut-off for inclusion of sequences in the profile set at e = 0.001) retrieved, with E <10-4, the sequences of NAIP, two uncharacterized human proteins and, unexpectedly, those of the mammalian telomerase-associated proteins (TP1) and a predicted NTPase from Streptomyces coelicolor. Further database searches initiated with these sequences showed significant similarity to two addnl. bacterial predicted NTPases (another one from Streptomyces coelicolor and one from Synechocystis sp.) and incompatibility locus protein from Podospora anserina (HET-E). The results of these searches suggested the existence of a new family of NTPases, which we termed the NACHT family, after NAIP, CIIA, HET-E and TP1. The multiple alignment of the NACHT proteins, constructed using the MACAW program, shows the conservation of seven distinct motifs, including the ATP/GTPase-specific P-loop, the Mg2+-binding site (Walker A and B motifs, resp.) and five more specific motifs.

REFERENCE COUNT:

THERE ARE 9 CITED REFERENCES AVAILABLE FOR 9

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L5 ANSWER 5 OF 47 SCISEARCH COPYRIGHT (c) 2005 The Thomson Corporation on

STN

ACCESSION NUMBER: 2003:1030234 SCISEARCH Full-text

THE GENUINE ARTICLE: 745DO

TITLE: Down-regulation of telomerase activity in malignant

glioma

cells by p27(KIP1)

AUTHOR:

Kanzawa T; Komata T; Kyo S; Germano I M; Kondo Y; Kondo

(Reprint)

CORPORATE SOURCE:

Univ Texas, MD Anderson Canc Ctr, Dept Neurosurg, 1515 Holcombe Blvd, Houston, TX 77030 USA (Reprint); Univ Texas, MD Anderson Canc Ctr, Dept Neurosurg, Houston, TX

77030 USA; Kanazawa Univ, Sch Med, Dept Obstet &

Gynecol,

Kanazawa, Ishikawa 9200934, Japan; CUNY Mt Sinai Sch

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Dept Neurosurg, New York, NY 10029 USA

COUNTRY OF AUTHOR:

USA; Japan

SOURCE:

INTERNATIONAL JOURNAL OF ONCOLOGY, (DEC 2003) Vol. 23,

No.

6, pp. 1703-1708.

Publisher: PROFESSOR D A SPANDIDOS, 1, S MERKOURI ST,

EDITORIAL OFFICE,, ATHENS 116 35, GREECE.

ISSN: 1019-6439.

DOCUMENT TYPE:

Article; Journal

LANGUAGE: English

REFERENCE COUNT: 46 *ABSTRACT IS AVAILABLE IN THE ALL AND IALL FORMATS*

AB The cyclin-dependent kinase inhibitor p27(KIP1) is considered not only a prognostic factor in cancer, but also a promising anti-cancer agent. However, the relationship between p27 KIP I and telomerase, that has potential as tumor-marker, remains to be elucidated. In this study, using the recombinant adenoviral vector expressing p27(KIP1) (Adp27(KIP1)), we investigated whether p27(KIP1) affects telomerase activity in malignant glioma U373-MG cells. Overexpression of p27(KIP1) suppressed telomerase activity in tumor cells. The downregulation of telomerase was due to inhibition of the-human telomerase reverse transcriptase (hTERT) gene expression at the transcriptional level. This inhibitory effect was partially induced by interfering with binding sites of the hTERT core promoter for transcription factors Myc and Spl. These findings identify a novel role for p27(KIP1) in down-regulation of telomerase activity.

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STN

ACCESSION NUMBER: 2000:526105 SCISEARCH Full-text

THE GENUINE ARTICLE: 332JN

TITLE: Identification and characterization of negative

regulatory elements of the human telomerase

catalytic subunit (hTERT) gene promoter: possible role

of

MZF-2 in transcriptional repression of hTERT

Fujimoto K; Kyo S (Reprint); Takakura M; Kanaya T; AUTHOR:

Kitagawa Y; Itoh H; Takahashi M; Inoue M

KANAZAWA UNIV, SCH MED, DEPT OBSTET & GYNECOL, KANAZAWA, CORPORATE SOURCE:

ISHIKAWA 920864, JAPAN (Reprint); KANAZAWA UNIV, SCH

MED,

DEPT OBSTET & GYNECOL, KANAZAWA, ISHIKAWA 920864, JAPAN; KANAZAWA UNIV, SCH MED, DEPT UROL, KANAZAWA, ISHIKAWA 920864, JAPAN; KANAZAWA UNIV, CANC RES INST, DEPT MOL &

CELLULAR BIOL, KANAZAWA, ISHIKAWA 920864, JAPAN

COUNTRY OF AUTHOR: **JAPAN**

SOURCE:

NUCLEIC ACIDS RESEARCH, (1 JUL 2000) Vol. 28, No. 13,

pp.

2557-2562.

Publisher: OXFORD UNIV PRESS, GREAT CLARENDON ST, OXFORD

OX2 6DP, ENGLAND. ISSN: 0305-1048.

DOCUMENT TYPE:

Article; Journal

FILE SEGMENT: LANGUAGE:

LIFE English

REFERENCE COUNT:

30

ABSTRACT IS AVAILABLE IN THE ALL AND IALL FORMATS AB

Human telomerase reverse transcriptase (hTERT) is a catalytic subunit of human telomerase and is a critical determinant of the enzymatic activity of telomerase. Expression of hTERT is known to be regulated mainly at the transcriptional level. In the present study, using transient expression assays, we identified a 400 bp silencer of the hTERT promoter between -776 and -378 upstream of the proximal core promoter. The inhibitory effects of this silencer were enhanced with

cellular differentiation. A computer-assisted homology search identified multiple binding motifs for myeloid-specific zinc finger protein 2 (MZF-2) within this region, Mutation introduced in these sites resulted in significant activation of hTERT transcription, Gel shift assays demonstrated that MZF-2 proteins specifically bound to these sites. Overexpression of MZF-2 in cells led to down-regulation of hTERT transcription as well as telomerase activity, These findings suggest that the 400 bp region upstream of the hTERT core promoter that we identified functions as a negative regulatory region and that MZF-2 may be an effector of negative regulation of hTERT.

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STN

ACCESSION NUMBER:

1998:945496 SCISEARCH Full-text

THE GENUINE ARTICLE: 146JK

TITLE:

Telomeres and telomerase: targets for cancer

chemotherapy?

AUTHOR:

Perry P J (Reprint); Kelland L R

CORPORATE SOURCE:

INST CANC RES, CANC RES CAMPAIGN BIOMOL STRUCT UNIT, 15 COTSWOLD RD, SUTTON SM2 5NG, SURREY, ENGLAND (Reprint);

INST CANC RES, CTR CANC THERAPEUT, SUTTON SM2 5NG,

SURREY,

ENGLAND

COUNTRY OF AUTHOR:

ENGLAND

SOURCE:

EXPERT OPINION ON THERAPEUTIC PATENTS, (DEC 1998) Vol.

8,

No. 12, pp. 1567-1586.

Publisher: ASHLEY PUBL LTD, 1ST FL, THE LIBRARY, 1 SHEPHERDS HILL HIGHGATE, LONDON N6 5QJ, ENGLAND.

ISSN: 1354-3776.

DOCUMENT TYPE:

General Review; Journal

FILE SEGMENT: LANGUAGE:

English 128

LIFE

REFERENCE COUNT:

ABSTRACT IS AVAILABLE IN THE ALL AND IALL FORMATS

AΒ Telomerase is a specialised ribonucleoprotein comprising of, at present, 3 known components: the human telomerase RNA component (hTR); the human telomerase reverse transcriptase catalytic subunit (hTERT), and TP1, a telomerase-associated protein. Applications involving telomerase have been proposed in the fields of cellular engineering, diagnostics/prognostics and therapeutics. In the diagnostics area, around 85% of human cancers have been shown to possess telomerase activity, while such activity is not detectable in most somatic cells. In some cases (notably neuroblastomas, gastric and breast rumours), higher levels of telomerase activity were associated with poor prognosis. Telomerase activity, which has generally been measured using a highly sensitive PCR-based TRAP assay, may also be assessed to monitor residual disease following surgery and/or chemotherapy. As telomerase appears to be selectively expressed in rumours versus normal cells, many have proposed that the enzyme represents a good target for inhibition. Efforts are underway to target various components of the telomerase/telomere machinery including the hTR template region using antisense oligonucleotides and peptide nucleic acids (PNAs), some of which inhibit at the nanomolar level, hTERT, and the telomere/telomerase interaction.

Small-molecule inhibitors of telomerase have recently been described. These include a series of regioisomeric diamidoanthracene-9,10-diones (the best of which inhibit telomerase in cell-free assays with IC50 values of 1 - 5 mu M) and porphyrin-based molecules. These molecules have been proposed to act via stabilisation of telomerase. Reverse transcriptase inhibitors, such as AZT triphosphate, and guanine-quadruplexes, structures associated with telomeres have also been shown to inhibit telomerase. This will clearly be an area where, in the near future, potent inhibitors will be developed thus permitting further target validation experiments to be performed in tumour-bearing mice and ultimately in cancer patients.

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on STN

ACCESSION NUMBER:

2003042499 EMBASE Full-text

TITLE:

Determinants in mammalian telomerase

RNA that mediate enzyme processivity and cross-species

incompatibility.

AUTHOR:

Chen J.-L.; Greider C.W.

CORPORATE SOURCE:

C.W. Greider, Department of Molecular Biology, Johns

Hopkins Univ. Sch. of Medicine, Baltimore, MD 21205,

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SOURCE:

EMBO Journal, (15 Jan 2003) Vol. 22, No. 2, pp. 304-314.

Refs: 39

ISSN: 0261-4189 CODEN: EMJODG

COUNTRY:

United Kingdom
Journal; Article

DOCUMENT TYPE: FILE SEGMENT:

029 Clinical Biochemistry

LANGUAGE:

English

SUMMARY LANGUAGE:

English

ENTRY DATE: En

Entered STN: 20030207 Last Updated on STN: 20030207

Telomerase contains two essential components: an RNA molecule that templates telomeric repeat synthesis and a catalytic protein component. Human telomerase is processive, while the mouse enzyme has much lower processivity. We have identified nucleotide determinants in the telomerase RNA that are responsible for this difference in processivity. Mutations adjacent to the template region of human and mouse telomerase RNA significantly altered telomerase processivity both in vitro and in vivo. We also identified functionally important nucleotides in the pseudoknot domain of telomerase RNA that potentially mediate the incompatibility between human TERT and mouse telomerase RNA. These experiments identify essential residues of the telomerase RNA that regulate telomerase activity and processivity.

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ACCESSION NUMBER: 2003:70360 SCISEARCH Full-text

THE GENUINE ARTICLE: 633FU

TITLE:

Down-regulation of telomerase activity via protein phosphatase 2A activation in salvicine-induced human

leukemia HL-60 cell apoptosis

AUTHOR: Liu W J; Jiang J F; Xiao D; Ding J (Reprint)

CORPORATE SOURCE: Chinese Acad Sci, Shanghai Inst Mat Med, Div Antitumor

Pharmacol, State Key Lab Drug Res, Shanghai Inst Biol

Sci,

Shanghai 200031, Peoples R China (Reprint)

COUNTRY OF AUTHOR: Peoples R China

SOURCE: BIOCHEMICAL PHARMACOLOGY, (15 DEC 2002) Vol. 64, No. 12,

pp. 1677-1687.

Publisher: PERGAMON-ELSEVIER SCIENCE LTD, THE BOULEVARD,

LANGFORD LANE, KIDLINGTON, OXFORD OX5 1GB, ENGLAND.

ISSN: 0006-2952.

DOCUMENT TYPE:

Article; Journal

LANGUAGE:

English

REFERENCE COUNT: 48

ABSTRACT IS AVAILABLE IN THE ALL AND IALL FORMATS AB Salvicine is a novel topoisomerase 11 inhibitor possessing significant antitumor activity, both in vitro and in vivo. The antitumor effect of salvicine is associated with its ability to induce tumor cell apoptosis. Telomerase plays an important role in the apoptotic pathway. However, little is known about the mechanisms of telomerase regulation during apoptosis induced by anticancer drugs. This study investigated the regulation of telomerase activity in salvicine-induced human leukemia HL-60 cell apoptosis. Salvicine treatment resulted in HL-60 cell apoptosis and down-regulation of telomerase activity in a time- and concentration-dependent manner. Repression of telomerase activity preceded a decrease in expression of the telomerase catalytic subunit (hTERT) and telomerase-associated protein (TP1) at the mRNA level, suggesting that the salvicineinduced decrease in telomerase activity may be additionally regulated by mechanisms other than telomerase subunit transcription. We observed that okadaic acid (OA), a protein phosphatase inhibitor, prevented the induction of apoptosis and the down-regulation of telomerase activity by salvicine. The significant increase in protein phosphatase 2A (PP2A) activity induced by salvicine treatment was blocked completely by OA. Moreover, although salvicine induced HL-60 cell apoptosis in a caspase-3-dependent manner, a specific caspase-3 inhibitor, Z-DEVD-FMK, did not prevent a decrease in telomerase activity or an increase in PP2A activity in apoptotic HL-60 cells, ruling out a role for caspase-3 in PP2A activation by salvicine. The results collectively suggest that the salvicine-induced decline in telomerase activity is not a consequence of HL-60 cell apoptosis and that it may be caused principally by the dephosphorylation of telomerase components mediated by PP2A activation. (C) 2002 Elsevier

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ACCESSION NUMBER: 2001:759870 SCISEARCH Full-text

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THE GENUINE ARTICLE: 473VR

TITLE: Telomerase reverse transcriptase and telomeric-repeat

binding factor protein 1 as regulators of

telomerase activity in pancreatic cancer cells

AUTHOR:

Yajima T; Yagihashi A; Kameshima H; Kobayashi D; Hirata

Κ;

Watanabe N (Reprint)

CORPORATE SOURCE: Sapporo Med Univ, Sch Med, Dept Clin Lab Med, Chuo Ku,

South 1, West 16, Sapporo, Hokkaido 0608543, Japan (Reprint); Sapporo Med Univ, Sch Med, Dept Clin Lab Med, Chuo Ku, Sapporo, Hokkaido 0608543, Japan; Sapporo Med Univ, Sch Med, Dept Surg, Sapporo, Hokkaido 0608543,

Japan

COUNTRY OF AUTHOR: Japan

SOURCE: BRITISH JOURNAL OF CANCER, (1 SEP 2001) Vol. 85, No. 5,

pp. 752-757.

Publisher: CHURCHILL LIVINGSTONE, JOURNAL PRODUCTION

DEPT,

ROBERT STEVENSON HOUSE, 1-3 BAXTERS PLACE, LEITH WALK,

EDINBURGH EH1 3AF, MIDLOTHIAN, SCOTLAND.

ISSN: 0007-0920.

DOCUMENT TYPE:

Article; Journal

LANGUAGE:

English

REFERENCE COUNT: 30

ABSTRACT IS AVAILABLE IN THE ALL AND IALL FORMATS,

AB Telomerase adds hexameric repeats of 5 ' -TTAGGG-3 ' termed telomeres to ends of chromosomal DNA. This enzyme has been implicated in cellular immortalization and cellular senescence. Recently. a number of relevant genes have been cloned, including these encoding three major components of human telomerase: human telomerase RNA component (hTR), human telomerase reverse transcriptase (hTERT), and telomerase-associated protein-1 (TEP1). Also important are genes encoding human telomeric-repeat binding factor protein (TRF) 1 and 2. To clarify mechanisms regulating telomerase activity, we studied telomerase activity, the telomeric restriction fragment (TRF) length and gene expression of these telomerase components and the telomericrepeat binding factor proteins in sequential observation following Xirradiation of cultured pancreatic cancer cells. We previously reported that PANC-1 cells are better able to tolerate thermal stress, antineoplastic drugs, and exposure to tumour necrosis factor than MIAPaCa-2 cells. MIAPaCa-2 and PANC-1 cells were exposed to Xirradiation, their telomerase activity was increased at 2 days and then decreased gradually. Of the three telomerase components, only hTERT mRNA expression showed parallel changes. TRF length was stable just before and after X-irradiation. Among binding factor proteins, TRF1 mRNA showed reciprocal changes possibly directed toward maintaining a stable telomere length. In this study, our results demonstrate that not only hTERT but also TRF1 are important regulator of telomerase activity. (C) 2001 Cancer Research Campaign.

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L1 4 S MAMMALIAN TELOMERASE AND (KIDNEY CELL LINE OR 293 CELL?)

L2 4 DUP REM L1 (0 DUPLICATES REMOVED)

L3 47 S MAMMALIAN TELOMERASE AND (INHIBITOR? OR ACTIVATOR? OR

REGULAT

L4 47 DUP REM L3 (0 DUPLICATES REMOVED)

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